Ree'd PCT/PTO 09 SEP 2004



10/506549

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/506,549

DATE: 09/09/2004 TIME: 16:15:50

Input Set : A:\SEQLIST 1361US.txt

Output Set: N:\CRF4\09092004\J506549.raw

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4 <110> APPLICANT: APPLERA CORPORATION
      6 <120> TITLE OF INVENTION: ISOLATED HUMAN TRANSPORTER PROTEINS,
             NUCLEIC ACID MOLECULES ENCODING HUMAN TRANSPORTER PROTEINS,
             AND USES THEREOF
     10 <130> FILE REFERENCE: CL001361-US
C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/506,549
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C--> 13 <141> CURRENT FILING DATE: 2004-09-03

15'<150> PRIOR APPLICATION NUMBER: 60/361,343

16 <151> PRIOR FILING DATE: 2002-03-05

18 <160> NUMBER OF SEQ ID NOS: 4

20 <170> SOFTWARE: FastSEQ for Windows Version 4.0

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24 <212> TYPE: DNA 25 <213 > ORGANISM: Homo sapiens 27 <400> SEQUENCE: 1 28 atgtcttcca agaagaatag aaagcggttg aaccaaagcg cggaaaatgg ttcgtccttg 60 29 ccctctgctg cttcctcttg tgcggaggca cgggctcctt ctgctggatc agacttcgcg 120 30 gcaacctccg ggactctgac ggtgaccaac ttattagaaa aggatgacaa aattcctaaa 180 31 acattccaga attcccttat tcatcttgga ctcaacacta tgaagtctgc aaatatatgt 240 32 ataggtcgac cagtgttgct tactagtttg aacggaaagc aagaggtgta tacagcctgg 300 33 cctatggcag gatttcctgg aggcaaggtc ggcctgagtg aaatggcaca gaaaaatgtg 360 34 ggtgtgaggc ctggtgatgc catccaggtc cagcctcttg tgggtgctgt gctacaggct 420 35 gaggaaatgg atgtggcact gagtgacaaa gatatggaaa ttaatgaaga agaactgact 480 36 ggttgtatcc tgagaaaact agatggcaag attgttttac caggcaactt tctgtattgt 540 37 acattctatg gacgaccgta caagctgcaa gtattgcgag tgaaaggggc agatggcatg 600 38 atattgggag ggcctcagag tgactctgac actgatgccc aaagaatggc ctttgaacag 660 39 tccagcatgg aaaccagtag cctggagtta tccttacagc taagccagtt agatctggag 720 40 gatacccaga teccaacate aagaagtact eettataaac caattgatga cagaattaca 780 41 aataaagcca gtgatgtttt gctggatgtt acacagagcc ctggagatgg cagtggactt 840 42 atgctagagg aagtcacagg tcttaaatgt aattttgaat ctgccagaga aggaaatgag 900 43 caacttactg aagaagagag actgctaaag ttcagcatag gagcaaagtg caatactgat 960 44 actttttatt ttatttcttc aacaacaaga gtcaatttta cagagattga taaaaattca 1020 45 aaaqaqcaaq acaaccaatt caaaqtaact tatgacatqa taggaggatt aagtagccag 1080 46 ctgaaagcaa ttagagaaat aattgaattg cccctcaaac agcctgagct tttcaagagt 1140 47 tatggaatte etgeeeetag aggagtgtta etttatggte etceaggtae tggaaaaaca 1200 48 atgategeea gggetgttge taatgaagtt ggageetatg tttetgtaat taatggteet 1260 49 gaaattataa gcaaattcta tggtgagact gaagcaaagt tacgtcagat atttgctgaa 1320 50 gccactctac gacacccatc aattattttt attgatgagc tggatgcact ttgtccgaaa 1380 51 agagaggggg cccagaatga agtggaaaaa agagttgtgg cttcactctt aacactgatg 1440 52 gatggcattg gttcagaagt aagtgaagga caagtgttgg ttcttggggc cacaaatcgc 1500 53 cctcatgcct tggatgctgc tctccgaaga cctgggcgat ttgataaaga gattgagatt 1560

54 ggagttccca atgctcagga ccggctagat attctccaga aactgcttcg aagggtaccc 1620

RAW SEQUENCE LISTINGPATENT APPLICATION: **US/10/506,549**DATE: 09/09/2004 TIME: 16:15:50

Input Set : A:\SEQLIST_1361US.txt
Output Set: N:\CRF4\09092004\J506549.raw

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57 aaacagccta acctccctga tgtcaaggtg gctggactgg tgaagattac tctgaaggat 1800
58 ttcttgcagg caatgaatga tatcagaccc agtgccatga gggaaatagc aattgatgtc 1860
59 ccaaatgtat cctggtcaga tataggagga ctggaaagta tcaaactgaa gttggaacag 1920
60 gctgtggaat ggcccttaaa acatccagag tctttcattc gaatgggtat tcagccacct 1980
61 aaaggagttc ttctctatgg gccacctggg tgctctaaaa caatgatagc aaaggctttg 2040
62 qccaatgaga gtggactgaa ttttctagct ataaaggggc ctgaattaat gaataaatat 2100
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64 tccattattt tctttgatga actggatgcc ttagcagttg aaaggggcag ttctttaggt 2220
65 gctgggaatg tagccgatcg tgttttggct cagctcttaa cagaaatgga tgggattgaa 2280
66 cagctaaagg atgtgaccat tttggcagct actaaccgtc cagataggat agacaaggct 2340
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68 agaagggaaa tatttaagct gcagtttcac tccatgcctg tcagtaatga agttgacctg 2460
69 gatgaactca teetteaaac egaegeatae teaggageag agattgtage tgtetgeaga 2520
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71 ttcactcagg ccttgagcac tgtgacacct agaattcctg agtcattgag acgtttttat 2640
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75 <211> LENGTH: 892
76 <212> TYPE: PRT
77 <213> ORGANISM: Homo sapiens
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82 Gly Ser Ser Leu Pro Ser Ala Ala Ser Ser Cys Ala Glu Ala Arq Ala
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84 Pro Ser Ala Gly Ser Asp Phe Ala Ala Thr Ser Gly Thr Leu Thr Val
85
86 Thr Asn Leu Leu Glu Lys Asp Asp Lys Ile Pro Lys Thr Phe Gln Asn
87
                           55
88 Ser Leu Ile His Leu Gly Leu Asn Thr Met Lys Ser Ala Asn Ile Cys
                       70
                                           75
90 Ile Gly Arg Pro Val Leu Leu Thr Ser Leu Asn Gly Lys Gln Glu Val
92 Tyr Thr Ala Trp Pro Met Ala Gly Phe Pro Gly Gly Lys Val Gly Leu
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               100
94 Ser Glu Met Ala Gln Lys Asn Val Gly Val Arg Pro Gly Asp Ala Ile
                               120
                                                   125
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96 Gln Val Gln Pro Leu Val Gly Ala Val Leu Gln Ala Glu Glu Met Asp
                           135
98 Val Ala Leu Ser Asp Lys Asp Met Glu Ile Asn Glu Glu Glu Leu Thr
                       150
100 Gly Cys Ile Leu Arg Lys Leu Asp Gly Lys Ile Val Leu Pro Gly Asn
101
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                    165
102 Phe Leu Tyr Cys Thr Phe Tyr Gly Arg Pro Tyr Lys Leu Gln Val Leu
                                    185
104 Arg Val Lys Gly Ala Asp Gly Met Ile Leu Gly Gly Pro Gln Ser Asp
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105
            195
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107		210	Com	T 0	~1	T 0	215	T	~1 -	T	0	220	T	3	T	a 1
		Ser	ser	ьец	GIU		Ser	ьец	GIII	ьeu		GIII	Leu	Asp	ьeu	
	225	mb ~	C1-	T1.	Dwa	230	C	70	Com	mb	235	m	T	D	T1.	240
	Asp	THE	GIII	тте		THE	Ser	Arg	ser		Pro	Tyr	гуѕ	Pro		Asp
111) an	7~~	Tla	Wh w	245	T	Ala	Com	7 an	250	T 011	T 011	7 ~~	77m 7	255	~1
113	Asp	Arg	116	260	ASII	пÃР	міа	ser	265	val	пеп	пеп	Asp		THE	GIII
	Sar	Dro	Glv		Cl v	Cor	Gly	T 011		T 011	C111	Cl.	17-1	270	C1	T 011
115	DCI	FIO	275	rsp	Gry		Gry	280	Mec	пец	Giu	Giu	285	1111	Gry	пец
	Lvs	Cvs		Phe	Glu	Ser	Ala		Glu	Glv	Δen	Glu		T.011	Thr	Glu
117		290	•••		014	001	295	**** 9	Ozu	019	11011	300	0111	пси	1111	Olu
			Ara	Leu	Leu	Lvs	Phe	Ser	Ile	Glv	Δla		Cvs	Asn	Thr	Asp
	305					310		501			315	_,5	Cyb	21011		320
		Phe	Tvr	Phe	Ile		Ser	Thr	Thr	Ara		Asn	Phe	Thr	Glu	
121			- 2 -		325					330					335	
122	Asp	Lys	Asn	Ser	Lys	Glu	Gln	Asp	Asn	Gln	Phe	Lvs	Val	Thr		Asp
123	-	•		340	•			-	345			•		350	•	
124	Met	Ile	Gly	Gly	Leu	Ser	Ser	Gln	Leu	Lys	Ala	Ile	Arg	Glu	Ile	Ile
125			355					360		_			365			
126	Glu	Leu	Pro	Leu	Lys	${\tt Gln}$	Pro	Glu	Leu	Phe	Lys	Ser	Tyr	Gly	Ile	Pro
127		370					375					380				
128	Ala	Pro	Arg	Gly	Val	Leu	Leu	Tyr	Gly	Pro	${\tt Pro}$	Gly	Thr	Gly	Lys	Thr
	385			,		390			•		395					400
130	Met	Ile	Ala	Arg	Ala	Val	Ala	Asn	Glu	Val	Gly	Ala	Tyr	Val	Ser	Val
131			•		405					410					415	•
	Ile	Asn	Gly		Glu	Ile	Ile	Ser	-	Phe	Tyr	Gly	Glu		Glu	Ala
133	_	_	_	420					425					430		
	Lys	Leu	_	GIn	Ile	Phe	Ala		Ala	Thr	Leu	Arg		Pro	Ser	Ile
135	- 1 -	D1	435	3	a 1	*		440	-	a		_	445	~3	~1	
	iie		TTE	Asp	GIU	ьеи	Asp	Ата	ьeu	Cys	Pro	_	Arg	GIU	GIA	Ala
137	~1 m	450	~1	77-7	~1	T	455	**~ 1	77-1	71-	C	460	7	m\	T	36
	465	ASII	GIU	vат	GIU	ьуs 470	Arg	vai	vaı	Ата	475	ьeu	Leu	Thr	Leu	мес 480
		Gl v	בוז	Glv	Sar		Val	Sor	G3 y	Glv		Tal	T 011	17-1	T 011	
141	тор	Gry	116	Gry	485	Giu	vai	per	Gru	490	GIII	vai	шец	vai	495	GIY
	Δla	Thr	Δen	Δra		Hic	Ala	T.e11	Δen		Δla	T.011	Ara	Δνα		Gl v
143	AΙα	1111	ASII	500	110	1110	AIG	пец	505	AIG	AIG	пец	Arg	510	PIO	GIY
	Ara	Phe	Asp		Glu	Tle	Glu	Tle		Val	Pro	Δen	Δla		Δen	Ara
145	9		515	-7.5				520	0 -1				525	· · · ·	1100	**** 9
	Leu	Asp		Leu	Gln	Lvs	Leu		Ara	Ara	Val	Pro		Leu	Leu	Thr
147		530				-1-	535		5	5		540				
	Glu		Glu	Leu	Leu	Gln	Leu	Ala	Asn	Ser	Ala		Glv	Tvr	Val	Glv
	545					550					555		2	- 4 -		560
		Asp	Leu	Lys	Val		Cys	Asn	Glu	Ala		Leu	Cys	Ala	Leu	
151		-		-	565		-			570	4		•		575	2
	Arg	Ile	Leu	Lys	Lys	Gln	Pro	Asn	Leu		Asp	Val	Lys	Val		Gly
153	_			580	_				585		-		-	590		-
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	Tro		Asp	Tle	Gly	Glv		Glu	Ser	Tle	Lvs		Lvs	Leu	Glu	Gln	
159	_				1	630					635		-1 -			640	
		Val	Glu	Trp	Pro	-	Lvs	His	Pro	Glu		Phe	Ile	Arq	Met		
161				F	645		-1-			650				5	655	2	
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163				660	-	-			665	-	•			670	-		
164	Lys	Thr	Met	Ile	Ala	Lys	Ala	Leu	Ala	Asn	Glu	Ser	Gly	Leu	Asn	Phe	
165	-		675			_		680					685				
166	Leu	Ala	Ile	Lys	Gly	Pro	Glu	Leu	Met	Asn	Lys	Tyr	Val	Gly	Glu	Ser	
167		690					695					700					
168	Glu	Arg	Ala	Val	Arg	Glu	Thr	Phe	Arg	Lys	Ala	Arg	Ala	Val	Ala	Pro	
169	705					710					715					720	
170	Ser	Ile	Ile	Phe	Phė	Asp	Glu	Leu	Asp	Ala	Leu	Ala	Val	Glu	Arg	Gly	
171					725					730					735		
172	Ser	Ser	Leu	Gly	Ala	Gly	Asn	Val	Ala	Asp	Arg	Val	Leu	Ala	Gln	Leu	
173				740					745					750			
174	Leu	Thr	Glu	Met	Asp	Gly	Ile	Glu	Gln	Leu	Lys	Asp	Val	Thr	Ile	Leu	
175			755					760					765				
176	Ala		Thr	Asn	Arg	Pro	-	Arg	Ile	Asp	Lys		Leu	Met	Arg	Pro	
177		770		_	_		775	_		_	_	780	_				
	_	Arg	Ite	Asp	Arg		He	Tyr	Val	Pro		Pro	Asp	Ala	Ala		
	785	3	a 1	- 1 -	Dl	790	T	a 1	Dl		795	1 7 - 1-	D	77-3	0	800	
	Arg	Arg	GIU	тте	Phe	гла	Leu	GIN	Pne		ser	Met	Pro	vai		Asn	
181	~1	77-7	7 ~~	T 011	805	~1	T 011	т1.	T 011	810	mb~	7	ח ד ת	m	815	C1	
183	GIU	vai	Asp	820	Asp	GIU	neu	116	825	GIII	1111	Asp	AIA	830	Ser	GIY	
	Λlo	Gl 11	т1Д		Ala	₩.	Cve	Δra		בוג	Δla	T.011	T.011		T.011	Gĺn	
185	Ата	Giu	835	vai	AIG	vai	Cys	840	Giu	АГа	лта	шец	845	AIG	Бец	GIU	
	Glu	Asp		Gln	Ala	Asn	Len		Met	Lvs	Ara	His		Thr	Gln	Ala	
187	014	850		02			855			_,,		860			0		
	Leu		Thr	Val	Thr	Pro		Ile	Pro	Glu	Ser		Ara	Arq	Phe	Tvr	
189				•		870		-		-	875			,		880	
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196	<212	2> TY	PE:	DNA													
197	97 <213> ORGANISM: Homo sapiens																
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	ccactcagtc						
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	taactctcag						
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	gggctcgaca						
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	agcatcgcgc						
	tccgttgccc						
	agagtgccat						
	ggagaccgcg						
	ggaaaacctc						
	gttggtttag						
	tacaatctaa						
	ctgtttcaaa						
237	tgcgctccac	tcaggggatt	acggcgcagg	ggcggaccct	cgctgacttc	tgccccggaa	1980
238	gtttttctct	cagttgaagc	gcgcacattg	agtcggcttt	tctactgctt	cggctagggt	2040
239	accttgtgac	catgtcttcc	aagaagaata	gaaagcggtt	gaaccaaagc	gcggaaaatg	2100
240	gttcgtcctt	gccctctgct	gcttcctctt	gtgcggaggc	acgggctcct	tctgctggat	2160
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	cagtaaaaat						
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RAW SEQUENCE LISTING ERROR SUMMARY DATE: 09/09/2004 PATENT APPLICATION: US/10/506,549 TIME: 16:15:51

Input Set : A:\SEQLIST_1361US.txt

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

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Seq#:3; N Pos. 77281,77282,77283,77284,77285,77286,77287,77288,77289,77290
Seq#:3; N Pos. 77291,77292,77293,77294,77295,77296,77297,77298,77299,77300
Seq#:3; N Pos. 77301,77302,77303,77304,77305,77306,77307,77308,77309,77310
Seq#:3; N Pos. 77311,77312,77313,77314,77315,77316,77317,77318,77319,77320
Seq#:3; N Pos. 77321,77322,77323,77324,77325,77326,77327,77328,77329,77330
Seq#:3; N Pos. 77331,77332,77333,77334,77335,77336,77337,77338,77339,77340
Seq#:3; N Pos. 77341,77342,77343,77344,77345,77346,77347,77348,77349,77350
Seq#:3; N Pos. 77351,77352,77353,77354,77355,77356,77357,77358,77359,77360
```

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 09/09/2004
PATENT APPLICATION: US/10/506,549 TIME: 16:15:51

Input Set : A:\SEQLIST 1361US.txt

Output Set: N:\CRF4\09092004\J506549.raw

Seq#:3; N Pos. 77361,77362,77363,77364,77365,77366,77367,77368,77369,77370
Seq#:3; N Pos. 77371,77372,77373,77374,77375,77376,77377,77378,77379,77380
Seq#:3; N Pos. 77381,77382,77383,77384,77385,77386,77387,77388,77389,77390
Seq#:3; N Pos. 77391,77392,77393,77394,77395,77396,77397,77398,77399,77400
Seq#:3; N Pos. 77401,77402,77403,77404,77405,77406,77407,77408,77409,77410
Seq#:3; N Pos. 77411,77412,77413,77414,77415,77416,77417,77418,77419,77420
Seq#:3; N Pos. 77421,77422,77423,77424,77425,77426,77427,77428,77429,77430

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/506,549

DATE: 09/09/2004

TIME: 16:15:51

Input Set : A:\SEQLIST 1361US.txt

Output Set: N:\CRF4\09092004\J506549.raw

L:12 M:270 C: Current Application Number differs, Replaced Current Application Number

L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:386 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:10860

M:341 Repeated in SeqNo=3